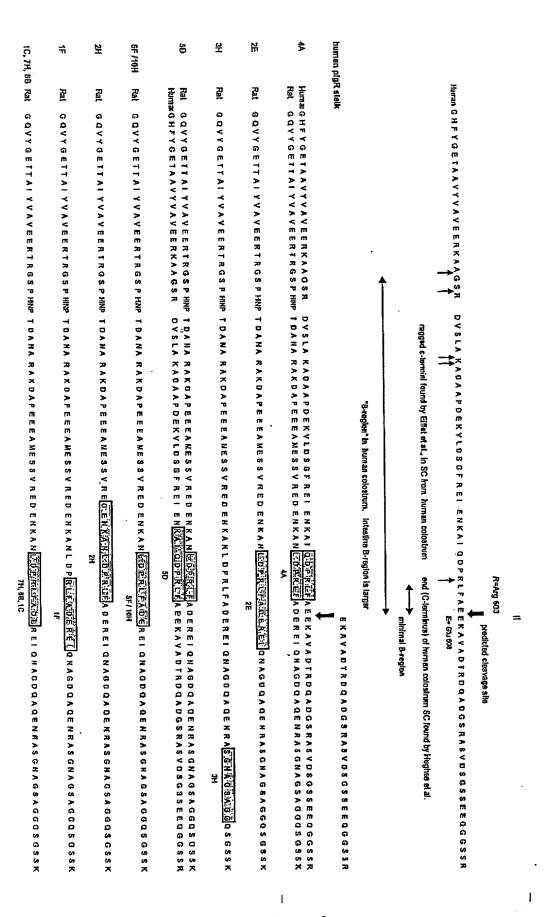
```
Ig-binding site
       Human
Bovine
Rat
Mouse
Possum
Rabbit
       VSSKYAGRANLTNFPENGTFVVNIAQLSQDDSGRYKCGLGINSRGLSFDVSLEVSQGPGLLNDTKVYTVDLGRTVTINC
Human
       Bovine
Rat
Mouse
Possum
Rabbit
       PFKTENAQKRKSLYKQIGLYPVLVIDSSGYV NPNYTGRIRLDIQGTGQLLFSVVINQLALSDAGQYLCQAGDDSNSN
--TRA-SE-----C-KTIQOCFQ-V--T--- SNS-KD-AHIS-L--NT-V-----RVX-----M-V-----AKAD
R--EG--HSK---C-KTRQSCE-----TEK---S-KD-AI-5MK--SRDI-Y-N-SH-IF----L-V---EGPSAD
--QEQ-T-D--F-C-KD-ESCA-----QEQ- G-D----A--S-S-SSRV-V-T-S-IXRQ-V-M-V-GV-S--DTGI
--TYATR-LK--F--VEDGEL--I----SKEAKD-R-K---T-Q--S-TAKE-T-T-KH-Q-N-----V--S-S-PTAE
Human
Bovine
Rat
Mouse
Possum
Rabbit
        Human
Bovine
Rat
Mouse
Possum
Rabbit
        369
Human
Bovine
Rat
Mouse
Possum
Rabbit
        445
Human
Bovine
Rat
muzeo9
Rabbit
                        Bridge to IgA
        Human
Bovine
Rat
Mouse
Possum
Rabbit
        Human
Bovine
Rat
Mouse
Possum
Rabbit
        ← Cytoplasmic
Human
Bovine
Mouse
 Possum
 Rabbit
                                                 Basolateral Targeting
                                                  Calmodulin Binding
        Human
 Bovine
 Mouse
 Rabbit
                                                    Rapid Endocytosis
                 → Avoid Degradation →
```

10 20 30 40 50 60 MLLFVLTCLL AVFPAISTKS PIFGPEEVNS VEGNSVSITC YYPPTSVNRH TRKYWCRQGA 70 80 90 100 110 120 RGGCITLISS EGYVSSKYAG RANLTNFPEN GTFVVNIAQL SQDDSGRYKC GLGINSRGLS 130 140 150 160 170 FDVSLEVSQG PGLLNDTXVY TVDLGRTVTI NCPFKTENAQ KRKSLYKQIG LYPVLVIDSS 200 210 220 230 240 GYVNPNYTGR IRLDIQGTGQ LLFSVVINQL RLSDAGQYLC QAGDDSNSNK KNADLQVLKP = 260 270 280 290 300 EPELVYEDLR GSVTFHCALG PEVANVAKFL CROSSGENCD VVVNTLGKRA PAFEGRILLN 320 330 340 350 PQDKDGSFSV VITGLRKEDA GRYLCGAHSD GQLQEGSPIQ AWQLFVNEES TIPRSPTVVK 370 380 390 400 410 GVAGSSVAVL CPYNRKESKS IKYWCLWEGA QNGRCPLLVD SEGWVKAQYE GRLSLLEEPG 440 450 460 470 NGTFTVILNQ LTSRDAGFYW CLTNGDTLWR TTVEIKIIEG EPNLKVPGNV TAVLGETLKV 500 510 520 530 PCHFPCKFSS YEKYWCKWNN TGCQALPSQD EGPSKAFVNC DENSRLVSLT LNLVTRADEG 550 - 560 570 580 590 -WYWCGVKQGH FYGETAAVYV AVEERKAAGS RDVSLAKADA APDEKVLDSG FREIENKAIQ 610 620 630 640 650 DPRLFAEEKA VADTRDQADG SRASVDSGSS EEQGGSSRAL VSTLVPLGLV LAVGAVAVGV 670 680 690 700 710 ARARHRONVD RVSIRSYRTD ISMSDFENSR EFGANDNMGA SSITQETSLG GKEEFVATTE 730 740 750 760 STTETKEPKK AKRSSKEEAE MAYKDFLLQS STVAAEAQDG PQEA

Figure 2

1



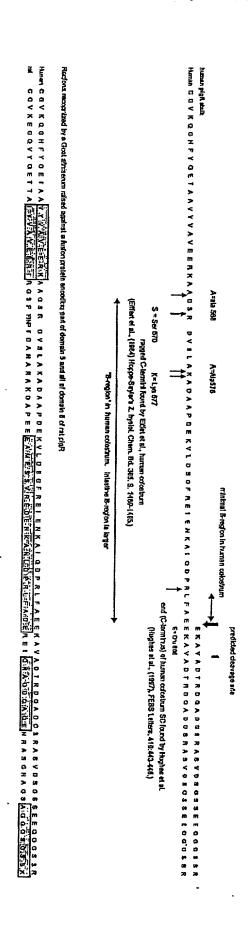


Figure 4

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FGQGTKVDIKRAAAEQKLISEEDINGAAHHHHHH	FR 2 WYQQKPGKAPKLLIY	CDR3 SFTVNSGYFQHWGQC	WVRQAPGKGLEWVSP	FR 2	MKYLLPTAAAGLLLL	PelB leader
mye 1QKL I S E E I	CDR 2 KASSLAS	FR 4 FTLVTVSS	ISGSGGS	CDR 2	AAQPAMA	
LNGAAHH	CDR 2 WYQQKPGKAPKLLIYKASSLASGVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQHYDSTPPT	CDR 3 FR 4 linker Light chain FR 1 CDR 1 SF'TVNSGYFQHWGQGTLVTVSSGGGGSGGGGSGGGSEIVLTQSPSTLSASIGDRVTITCRASEGIYHWLA	TYYADSVK	R 2	DYKAKQVQ	FLAG
НННН вня			WVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAR	FR 3	MKYLLPTAAAGLLLLAAQPAMADYKAKQVQLVQSGGGLVQPGGSLRLSCAASGFTFSSYAMS	Heavy chain FR 1
					TFSSYAMS	CDR 1
Ĺ	-	CDR 1 SEGIYHWLA				

determining regions (CDR) of the heavy chain and light chain are indicated. repeated three times), a light chain variable region, a myc epitope tag and a 6HIS tag (for purification by (for secretion in E. coli), a FLAG epitope tag, a heavy chain variable region, a linker sequence (GGGS The amino acid sequence of the secreted form of the ScFv 4AF is shown. The ScFv consists of a pelb leader Immobilized Metal-ion Affinity Chromatography (IMAC)). The framework (FR) and complementarity-